

SUPPLEMENTARY TABLE S2. CLASSIFICATION OF EXPRESSION QUANTITATIVE TRAIT LOCUS STUDIES BY CELL TYPE AND POPULATION

<i>Class</i>	<i>Tissue</i>	<i>Population</i>	<i>Studies</i>	<i>Datasets</i>
AllCell_AllPop	All	All	16	All
LCL_CEU	LCL	CEU	8	HA_CEU, HA2_CEU, HRC, AS, MRC, 3CL, EGEUV_EUR, MuTHER_LCL
LCL_YRI	LCL	YRI	4	HA_YRI, HA2_YRI, HRY, EGEUV_YRI
LCL_ASN	LCL	ASN	1	HA_CHB, HA_JPT
LCL	LCL	All	9	HA_CEU, HA_YRI, HA_CHB, HA_JPT, HA2_CEU, HA2_YRI, HRC, HRY, AS, MRC, 3CL, EGEUV_EUR, EGEUV_YRI, MuTHER_LCL
Bcell	B cell	All	1	IM_B
Monocyte	Monocyte	All	2	MO, IM_MO
Tcell	T cell	All	1	3CT
Brain	Brain	All	2	BR, BR2_Cer, BR2_FC, BR2_TC, BR2_P
Liver	Liver	All	2	LV, LV2
Skin	Skin	All	2	SKN, MuTHER_Skin
Fibroblast	Fibroblast	All	1	3CF
Fat	Adipose	All	1	MuTHER_Fat

ASN indicates pooled CHB+JPT populations.
LCL, lymphoblastoid cell line.